AMENDMENT

In the specification:

Please replace table 5, beginning at page 31, with the rewritten table 5:

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AA	Sequence	SEQ ID NO	Antigen	Protein or Molecule	1 st Position	A*0201
8	ALPPVAPV	107	P53		69	0.0230
11	LLPENNVLSPV	108	P53		25	0.1300
9	SLYNTITVL	109	HIV	gag	77	0.0330
9	SLYNTISVL	110	HIV	gag	77	0.0190
9	SLYNTVSTL	111	HIV	gag	77	0.0320
9	AIYGRPVSA	112	KSHV		508	0.0560
9	ALIGTMCGI	113			237	0.1500
9	ATLGTVILL	114	KSHV		8	0.0280
9	FIALNLSFI	115	KSHV		624	0.0640
9	FIQNIDFKA	116	KSHV		631	0.1400
9	FLNSSNLFT	117	KSHV		560	0.0780
9	FLYVVCSLA	118	KSHV		2	1.1000
9	FVAVHVPDV	119	KSHV		8	0.2600
9	GILGTIIFA	120	KSHV		244	0.0270
9	HLDFWHHEV	121	KSHV		168	0.2400
9	ITATFTAPL	122	KSHV		342	0.1300
9	LLGTWMFSV	123	KSHV		53	1.5000
9	LMWYELSKI	124	KSHV		492	0.0670
9	MIIIVIAII	125	KSHV	•	738	0.0150
9	NLLDRLLLI	126	KSHV		77	0.0290
9	RIFYNILEI	127	KSHV		20	0.0800
9	RLASSVFDL	128	KSHV		649	0.0670
9	RLGAIPPLV	129	KSHV		24	0.0150
	8 11 9 9 9 9 9 9 9 9 9 9 9 9	8 ALPPVAPV 11 LLPENNVLSPV 9 SLYNTITVL 9 SLYNTISVL 9 SLYNTVSTL 9 AIYGRPVSA 9 ALIGTMCGI 9 ATLGTVILL 9 FIALNLSFI 9 FIQNIDFKA 9 FLNSSNLFT 9 FLYVVCSLA 9 FVAVHVPDV 9 GILGTIIFA 9 HLDFWHHEV 9 ITATFTAPL 9 LLGTWMFSV 9 LMWYELSKI 9 MIIIVIAII 9 NLLDRLLLI 9 RIFYNILEI 9 RLASSVFDL	NO 8 ALPPVAPV 107 11 LLPENNVLSPV 108 9 SLYNTITVL 109 9 SLYNTISVL 110 9 SLYNTVSTL 111 9 AIYGRPVSA 112 9 ALIGTMCGI 113 9 ATLGTVILL 114 9 FIALNLSFI 115 9 FIQNIDFKA 116 9 FLNSSNLFT 117 9 FLYVVCSLA 118 9 FVAVHVPDV 119 9 GILGTIIFA 120 9 HLDFWHHEV 121 9 ITATFTAPL 122 9 LLGTWMFSV 123 9 LMWYELSKI 124 9 MIIIVIAII 125 9 NLLDRLLLI 126 9 RIFYNILEI 127 9 RLASSVFDL 128	NO 8 ALPPVAPV 107 P53 11 LLPENNVLSPV 108 P53 9 SLYNTITVL 109 HIV 9 SLYNTISVL 110 HIV 9 SLYNTVSTL 111 HIV 9 SLYNTVSTL 111 HIV 9 AIGTMCGI 113 SHV 9 ALIGTWCGI 113 SHV 9 FIALNLSFI 115 KSHV 9 FIQNIDFKA 116 KSHV 9 FLYVVCSLA 118 KSHV 9 FLYVVCSLA 118 KSHV 9 FVAVHVPDV 119 KSHV 9 GILGTIIFA 120 KSHV 9 HLDFWHHEV 121 KSHV 9 LLGTWMFSV 123 KSHV 9 MIIIVIAII 125 KSHV 9 NLLDRLLLI 126 KSHV 9 RIFYNILEI 1	NO Molecule 8 ALPPVAPV 107 P53 11 LLPENNVLSPV 108 P53 9 SLYNTITVL 109 HIV gag 9 SLYNTISVL 110 HIV gag 9 SLYNTVSTL 111 HIV gag 9 SLYNTVSTL 111 HIV gag 9 AIYGRPVSA 112 KSHV 9 ALIGTMCGI 113 SHV 9 FIALNLSFI 115 KSHV 9 FIQNIDFKA 116 KSHV 9 FLYVVCSLA 118 KSHV 9 FVAVHVPDV 119 KSHV 9 FVAVHVPDV 119 KSHV 9 HLDFWHHEV 121 KSHV 9 ITATFTAPL 122 KSHV 9 LLGTWMFSV 123 KSHV 9 MIIIVIAII 125 KSHV 9 RIFYNILEI <	NO Molecule Position 8 ALPPVAPV 107 P53 69 11 LLPENNVLSPV 108 P53 25 9 SLYNTIVL 109 HIV gag 77 9 SLYNTVSTL 110 HIV gag 77 9 SLYNTVSTL 111 HIV gag 77 9 AIYGRPVSA 112 KSHV 508 9 ALIGTMCGI 113 237 9 ATLGTVILL 114 KSHV 8 9 FIQNIDFKA 116 KSHV 624 9 FLNSSNLFT 117 KSHV 560 9 FLYVVCSLA 118 KSHV 2 9 FLYVVCSLA 118 KSHV 244 9 HLDFWHHEV 121 KSHV 244 9 HLDFWHHEV 121 KSHV 342 9 LLGTWMFSV 123 KSHV 492

Peptide	AA	Sequence	SEQ ID NO	Antigen	Protein or Molecule	1 st Position	A*0201
32.0078	9	RLYQASAVM	130	KSHV		4	0.0180
32.0081	9	SILGCDVSV	131	KSHV		226	0.0430
32.0087	9	SVDFYQFRV	132	KSHV		59	0.0160
32.0088	9	SVSDFDLRI	133	KSHV	•	245	0.0120
32.0090	9	TLGTVILLV	134	KSHV		9	0.0830
32.0099	9	YLVWQPMSA	135	KSHV		398	0.0130
32.0114	10	AAVEQILTSV	136	KSHV		237	0.0210
32.0118	10	ALIGTMCGIL	137	KSHV		237	0.0120
32.0120	10	ATLGTVILLV	138	KSHV		8	0.0690
32.0124	10	FLYVVCSLAV	139	KSHV		2	0.2400
32.0127	10	GALPICSFVV	140	KSHV		27	0.0160
32.0137	10	KLLGTWMFSV	141	KSHV		52	1.6000
32.0148	10	QLASILGCDV	142	KSHV		223	0.0160
32.0150	10	RLSNRICFWA	143	KSHV		164	0.0130
32.0154	10	SLVTGFINFI	144	KSHV		720	0.0210
32.0159	10	VLATDVTSFL	145	KSHV		149	0.0190
32.0160	10	VLLNGWRWRL	146	KSHV		16	0.2800
32.0164	10	YLVWQPMSAI	147	KSHV		398	0.0230
34.0006	8	QLAKTCPV	148	P53		136	0.0110
34.0132	9	ALBRWGLLV	149	HER2/ neu		5	0.0960
34.0133	9	ALCRWGLLV	150	HER2/ neu		5	0.0360
34.0134	9	AMCRWGLLV	151	HER2/ neu		5	0.0280
34.0135	9	KLFGSLAFL	152	HER2/ neu		369	0.1200
34.0136	9	KLFGSLAFV	153	HER2/ neu		369	0.0900
34.0137	9	KMFGSLAFL	154	HER2/ neu		369	0.2000
34.0138	9	KMFGSLAFV	155	HER2/ neu		369	0.1200

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Peptide	AA	Sequence	SEQ ID NO	Antigen	Protein or Molecule	1 st Position	A*0201
34.0140	9	KMFBQLAKV	156	P53		132	0.0980
34.0141	9	KMFCQLAKV	157	p53		132	0.0400
34.0199	10	RMPEAAPPVV	158	p53		65	0.0340
34.0201	10	ALNKMFCQLV	159	p53		129	0.0240
35.0031	9	VLLVSLGAI	160	Flu	HEMA	542	0.0170
35.0034	9	LLTEVETPI	161	Flu	VMT2	3	0.2100
35.0036	9	RLIQNSLTI	162	Flu	VNUC	55	0.0300
35.0040	9	KMNIQFTAV	163	Flu	HEMA	402	0.0330
35.0046	10	GLFGAIAGFI	164	Flu	HEMA	345	0.0451
35.0047	10	KLESMGIYQI	165	Flu	HEMA	521	0.0120
35.0048	10	SLPFQNIHPV	166	Flu	HEMA	306	0.0520
37.0013	8	ALPPVAPV	167	p53		69	0.0500
37.0015	9	ALNKMFCQV	168	p53		129	0.0770
37.0017	9	KLFCQLAKV	169	p53		132	0.0640
37.0018	9	KLCPVQLWV	170	p53		139	0.0440
37.0019	9	CLTIHYNYV	171	p53		229	0.0110
37.0032	10	ALNKMFCOLV	172	p53		129	0.0150
37.0033	10	VLVPYEPPEV	173	p53		216	0.1100
37.0034	10	RLPEAAPPVV	174	p53		65	0.0350
37.0035	10	LLPPQHLIRV	175	p53		188	0.0120
37.0069	11	ILLEDSSGNLV	176	p53		255	0.0590
F124.03	10	KLVALGINAV	177	HCV	NS3	1406	0.0110
F124.04	9	SLMAFTAAV	178	HCV	NS4	1789	0.1900
F124.06	9	CINGVCWTV	179	HCV	NS3	1073	0.0910
F124.08	9	TISGVLWQV	180	HCV	NS3	1073	0.1400
F124.09	9	SISGVLWQV	181	HCV	NS3	1073	0.1400
F124.10	9	SLMAFTASV	182	HCV	NS4	1789	0.1200
F124.11	9	GLRDCTMLV	183	HCV	NS5	2727	0.0120
F124.12	10	KLVALGVNAV	184	HCV	NS3	1406	0.0200
F124.14	10	KLSGLGLNAV	185	HCV	NS3	1406	0.0170
F124.23	10	KLVSLGVNAV	186	HCV	NS3	1406	0.0150
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